



Fig 1A

5'-UTS →
5-GACCCTAGAT CCAAGATCT CCAAGGATTT GGTGGCATAC CCACTCCAGC ACACAGAAAGC ATGAGGTTCA
start codon
exon 1 intron 1 exon 2
TGACTCTCCT CTTCCTGACA GCTCTGGCAG GAGCCCTGGT CTGTGCCT ↑ ATGATCCAGAGG CCGCCTCTGC
intron 2 exon 3
CCCAGGATCG GGGAAACC ↑ CTTGCCATGAAGC ATCAGCAGCT CAAAAGGAAA ATGCAGGTGA AGACCCAGGG
intron 3 exon 4
TTAGCCAGAC AGGCACCAA GCCAAGGAAG CAGAGATCCA GCCTTCTGG ↑ A AAAAGGCCTA GACGGAGCAA
intron 4
AAAAAGCTGT GGGGGGACTC GGAAAACTAG GAAAAGATGC AGTCGAAGAT CTAGAAAAGCG TGGGTAAAGG ↑
exon 5 stop codon
AGCCGTCCAT GACGTTAAAG ACGTCCTTGA CTCAGTACTA TAGCTGTAAAG GAGAAAGCTGA GAAATGATAC
← 3'-UTS
CCAGGAGCAG CAGGCTTTAC GTTTTCAGCC TAAACCT

Fig 1B

MRFMILLFLT ALAGALVCAY DPEAASAPGS GNPCHASAA QKENAGEDPG LARQAPKPRK QRSSLEKGL
DGAKKAVGGL GKLGKDAVED LESVGKGAVH DVKDVLDVSL

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DCD-eGFP on various microbial pathogens

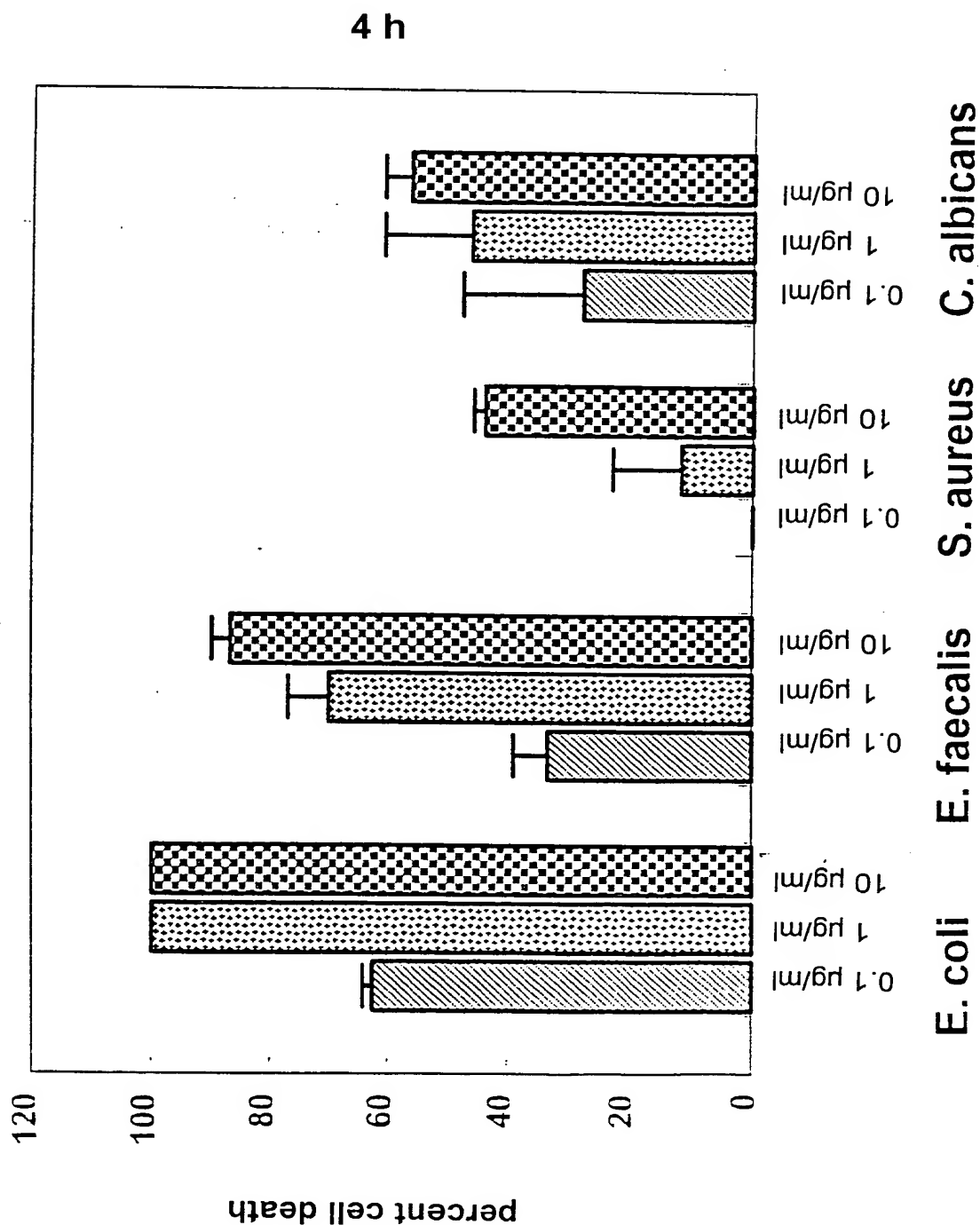


Fig. 2

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peptide of SEQ-ID No. 2 on various microbial pathogens

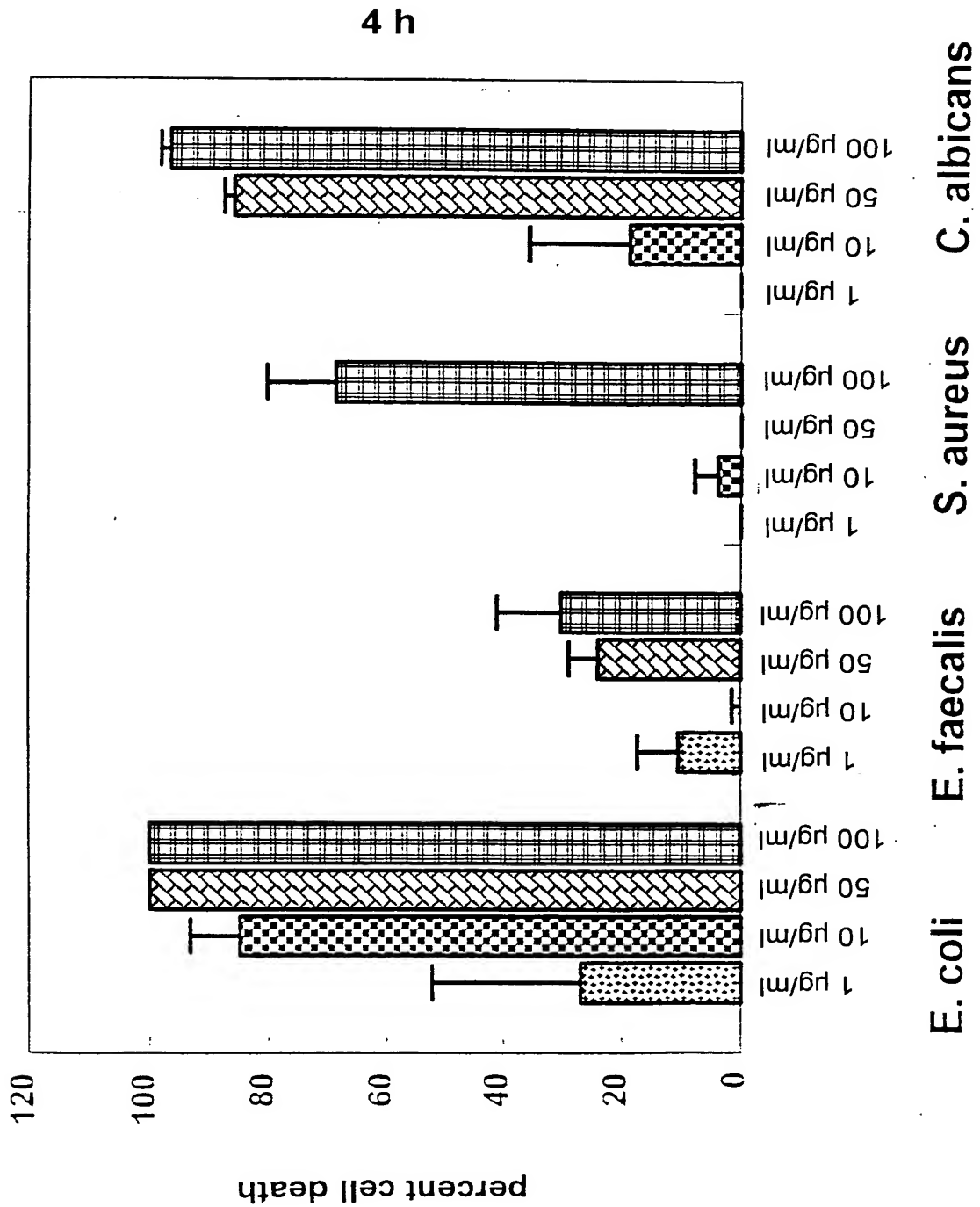
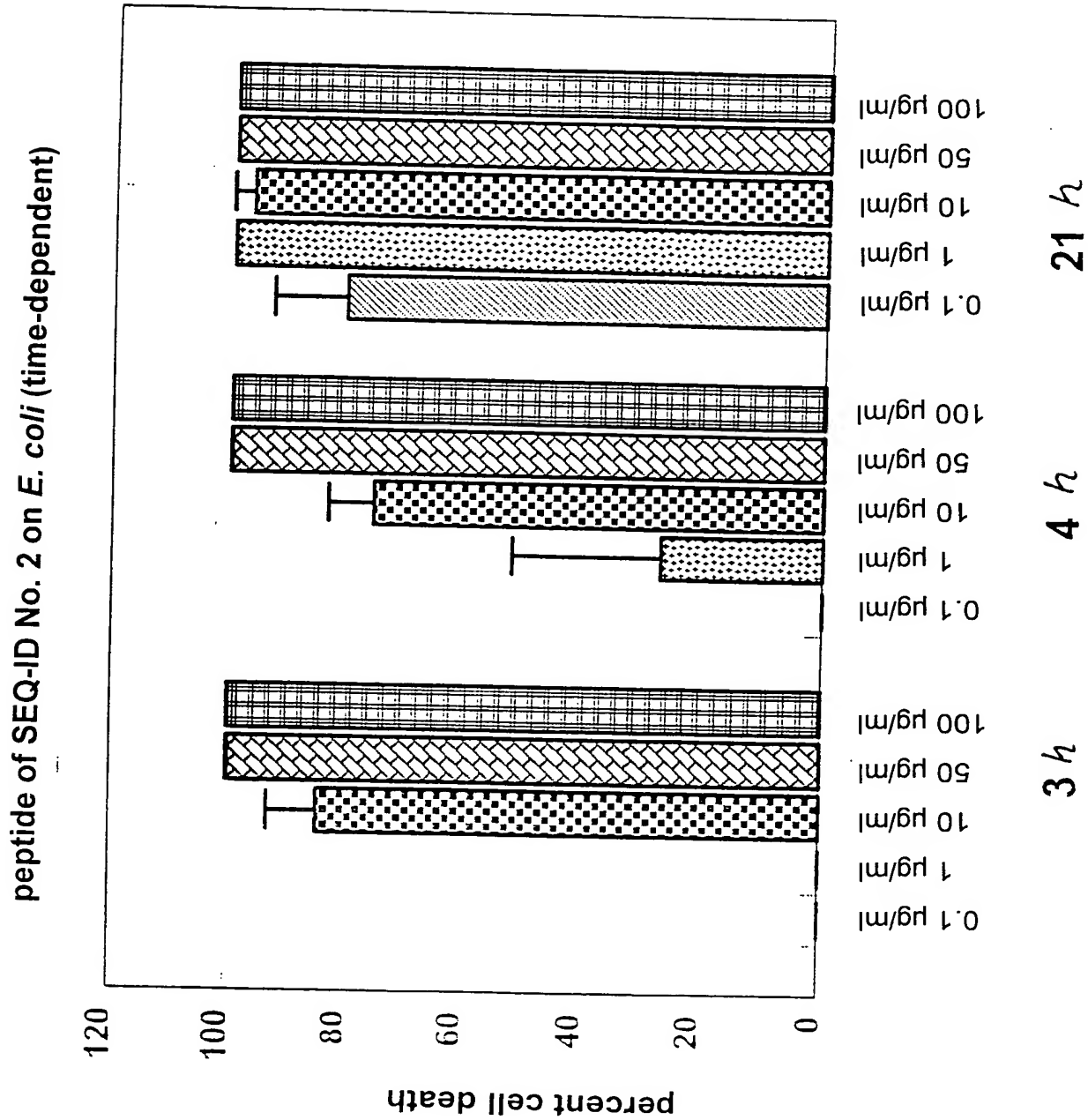


Fig. 3

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Fig. 4



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peptide of SEQ ID No. 3 on various microbial pathogens

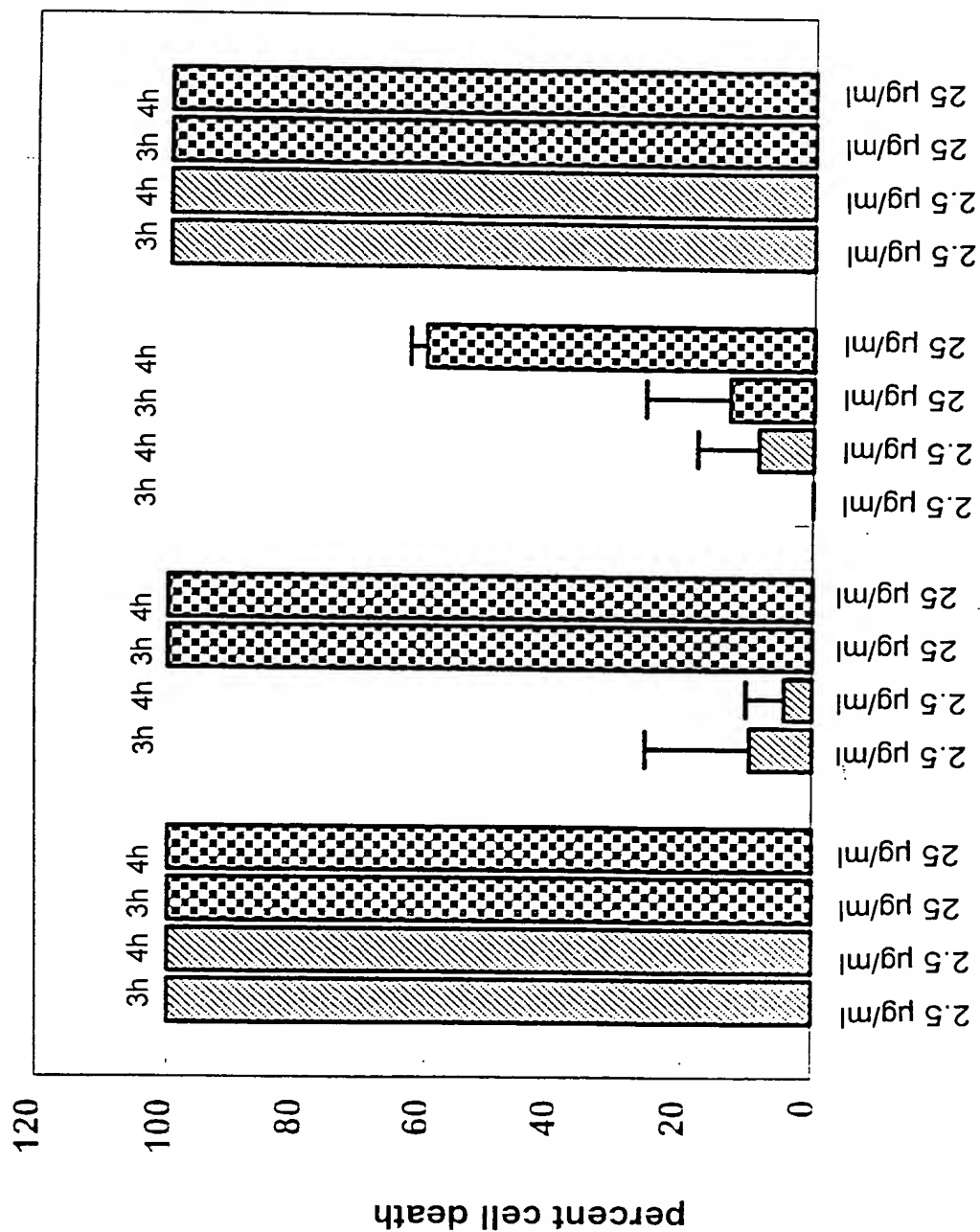


Fig. 5